



SEQUENCE LISTING

RECEIVED

NOV 16 2000

TECH CENTER 1600/2800

<110> Cho, Myeong G.
Lemaux, Peggy G.
Buchanan, Bob B.
Wong, Joshua
Marx, Corina

<120> Value-Added Traits in Grain and Seed
Transformed with Thioredoxin

<130> 2001-0703.30

<140> US 09/538,864

<141> 2000-03-29

<150> US 60/126,736

<151> 1999-03-29

<150> US 60/127,198

<151> 1999-03-31

<150> US 60/169,162

<151> 1999-12-06

<150> US 60/177,740

<151> 2000-01-21

<150> US 60/177,739

<151> 2000-01-21

<160> 25

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 486

<212> DNA

<213> Artificial Sequence

<220>

<223> barley B1-hordein promoter and signal sequence

<400> 1

| | | | | | | |
|------------|------------|-------------|------------|-------------|------------|-----|
| aagctttaac | aaccacaca | ttgattgcaa | cttagtccta | cacaagtttt | ccattcttgt | 60 |
| ttcaggctaa | caacctatac | aaggttccaa | aatcatgcaa | aagtgatgct | aggttgataa | 120 |
| tgtgtgacat | gtaaagtga | taagggtgagt | catgcatacc | aaacctcggg | atttctatac | 180 |
| tttgtgtatg | atcatatgca | caactaaaag | gcaactttga | ttatcaattg | aaaagtaccg | 240 |
| cttgtagctt | gtgcaacct | acacaatgtc | caaaaatcca | tttgcaaaaag | catccaaaca | 300 |
| caattgttaa | agctgttcaa | acaaacaaaag | aagagatgaa | gcctggctac | tataaatagg | 360 |
| caggtagtat | agagatctac | acaagcacia | gcatcaaaac | caagaaacac | tagttaaac | 420 |
| caatccacta | tgaagacctt | cctcatcttt | gcactcctcg | ccattgcggc | aacaagtacg | 480 |
| attgca | | | | | | 486 |

<210> 2

<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> barley B1-hordein signal protein

<400> 2
Met Lys Thr Phe Leu Ile Phe Ala Leu Leu Ala Ile Ala Ala Thr Ser
1 5 10 15
Thr Ile Ala

<210> 3
<211> 497
<212> DNA
<213> Artificial Sequence

<220>
<223> Barley D-hordein promoter and signal sequence

<400> 3
cttcgagtgcc cgcgcgattt gccagcaatg gctaacagac acatattctg ccaaaacccc 60
agaacaataa tcacttctcg tagatgaaga gaacagacca agatacaaac gtccacgctt 120
cagcaaacag taccacagaa ctaggattaa gccgattacg cggcttttagc agaccgtcca 180
aaaaaactgt ttgcaaagc tccaattcct ccttgcttat ccaatttctt ttgtgttggc 240
aaactgcact tgtccaaccg attttggtct tcccggtgtt cttcttaggc taactaacac 300
agccgtgcac atagccatgg tccggaatct tcacctcgtc cctataaaag ccagccaat 360
ctccacaatc tcatcatcac cgagaacacc gagaaccaca aaactagaga tcaattcatt 420
gacagtccac cgagatggct aagcggctgg tcctctttgt ggcggtaatc gtgcgcctcg 480
tggtctcac caccgct 497

<210> 4
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> barley D-hordein signal protein

<400> 4
Ala Lys Arg Leu Val Leu Phe Val Ala Val Ile Val Ala Leu Val Ala
1 5 10 15
Leu Thr Thr Ala
20

<210> 5
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 5
atatctagaa tggcggcgtc ggcggcga 28

<210> 6
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 6
 atagagctct tactgggccg cgtgtag 27

 <210> 7
 <211> 26
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 7
 gtaaagcttt aacaacccac acattg 26

 <210> 8
 <211> 34
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 8
 ccgacgccgc tgcaatcgta cttgttgccg caat 34

 <210> 9
 <211> 33
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 9
 agaaagcttg gtacccttcg agtgcccgcc gat 33

 <210> 10
 <211> 48
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 10
 gaacagctcc tcgcccttgc tcacagcggg ggtgagagcc acgagggc 48

 <210> 11
 <211> 19

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 11
 ccaagaagtt cccagctgc 19

 <210> 12
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 12
 aactctagac tcggtggact gtcaatg 27

 <210> 13
 <211> 25
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 13
 catcgagaca agcacggtca acttc 25

 <210> 14
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 14
 atatccgagc gcctcgtgca tgcg 24

 <210> 15
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 15
 caagatggat tgcacgcagg ttct 24

 <210> 16
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>

<223> primer

<400> 16

atagaaggcg atgcgctgcg aat

23

<210> 17

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 17

cggaattcga tctagtaaca tagatgaca

29

<210> 18

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 18

ggtctagaat ggaaactcac aaaacc

26

<210> 19

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 19

atagctgcga caaccctgtc ott

23

<210> 20

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 20

gggagctctc aatcactctt accctc

26

<210> 21

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 21

aagcctgaac tcaccgac g

21

<210> 22

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 22

aagaccaatg cggagcatat ac

22

<210> 23

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> primer

<400> 23

ggcgcatgcy aattcgaatt cgatatcgat cttega

36

<210> 24

<211> 369

<212> DNA

<213> barley

<220>

<221> misc_feature

<222> (0)...(0)

<223> thioredoxin h

<400> 24

| | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| atggcgcggt | cggcaacggc | ggcggcagtg | gcggcgagg | tgatctcggt | ccacagcctg | 60 |
| gagcagtgga | ccatgcagat | cgaggaggcc | aacaccgcca | agaagctggt | ggtgattgac | 120 |
| ttcactgcat | catggtgagg | accatgccgc | atcatggctc | cagttttcgc | tgatctcgcc | 180 |
| aagaagttcc | caaagtctgt | tttcctcaag | gtcgacgtgg | atgaactgaa | gcccattgct | 240 |
| gagcaattca | gtgtcgaggc | catgccaaag | ttcctgttca | tgaaggaagg | agacgtcaag | 300 |
| gacagggttg | tcggagctat | caaggaggaa | ctgaccgcca | aggttgggct | tcaocgggcg | 360 |
| gcccagtaa | | | | | | 369 |

<210> 25

<211> 122

<212> PRT

<213> barley

<220>

<221> VARIANT

<222> (0)...(0)

<223> thioredoxin h

<400> 25

| | | | | | | | | | | | | | | | |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Met | Ala | Ala | Ser | Ala | Thr | Ala | Ala | Ala | Val | Ala | Ala | Glu | Val | Ile | Ser |
| 1 | | | | 5 | | | | | 10 | | | | | 15 | |
| Val | His | Ser | Leu | Glu | Gln | Trp | Thr | Met | Gln | Ile | Glu | Glu | Ala | Asn | Thr |
| | | | 20 | | | | | 25 | | | | | 30 | | |
| Ala | Lys | Lys | Leu | Val | Val | Ile | Asp | Phe | Thr | Ala | Ser | Trp | Cys | Gly | Pro |
| | | 35 | | | | | 40 | | | | | 45 | | | |
| Cys | Arg | Ile | Met | Ala | Pro | Val | Phe | Ala | Asp | Leu | Ala | Lys | Lys | Phe | Pro |
| | 50 | | | | | 55 | | | | | 60 | | | | |
| Asn | Ala | Val | Phe | Leu | Lys | Val | Asp | Val | Asp | Glu | Leu | Lys | Pro | Ile | Ala |
| 65 | | | | | 70 | | | | | 75 | | | | 80 | |
| Glu | Gln | Phe | Ser | Val | Glu | Ala | Met | Pro | Thr | Phe | Leu | Phe | Met | Lys | Glu |
| | | | 85 | | | | | | 90 | | | | | 95 | |
| Gly | Asp | Val | Lys | Asp | Arg | Val | Val | Gly | Ala | Ile | Lys | Glu | Glu | Leu | Thr |
| | | | 100 | | | | | 105 | | | | | 110 | | |
| Ala | Lys | Val | Gly | Leu | His | Ala | Ala | Ala | Gln | | | | | | |
| | | 115 | | | | | | 120 | | | | | | | |